

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage YungJamal complete sequence, 70214 bp including 4-base 3' overhang (GTC  
Analysis Date : 3/17/18 at 16:24:50  
Pages : 35  
Sequence Length : 70214 bp  
GC Content : 65.34%

Window Length : 96 bp  
Window Step : 12 bp  
Threshold Value : 0.500

PS-Version : 1.2

GeneMark Options : PostScript graph,  
Mark ORFs / splice sites,  
List ORFs,  
List regions and/or splice sites,

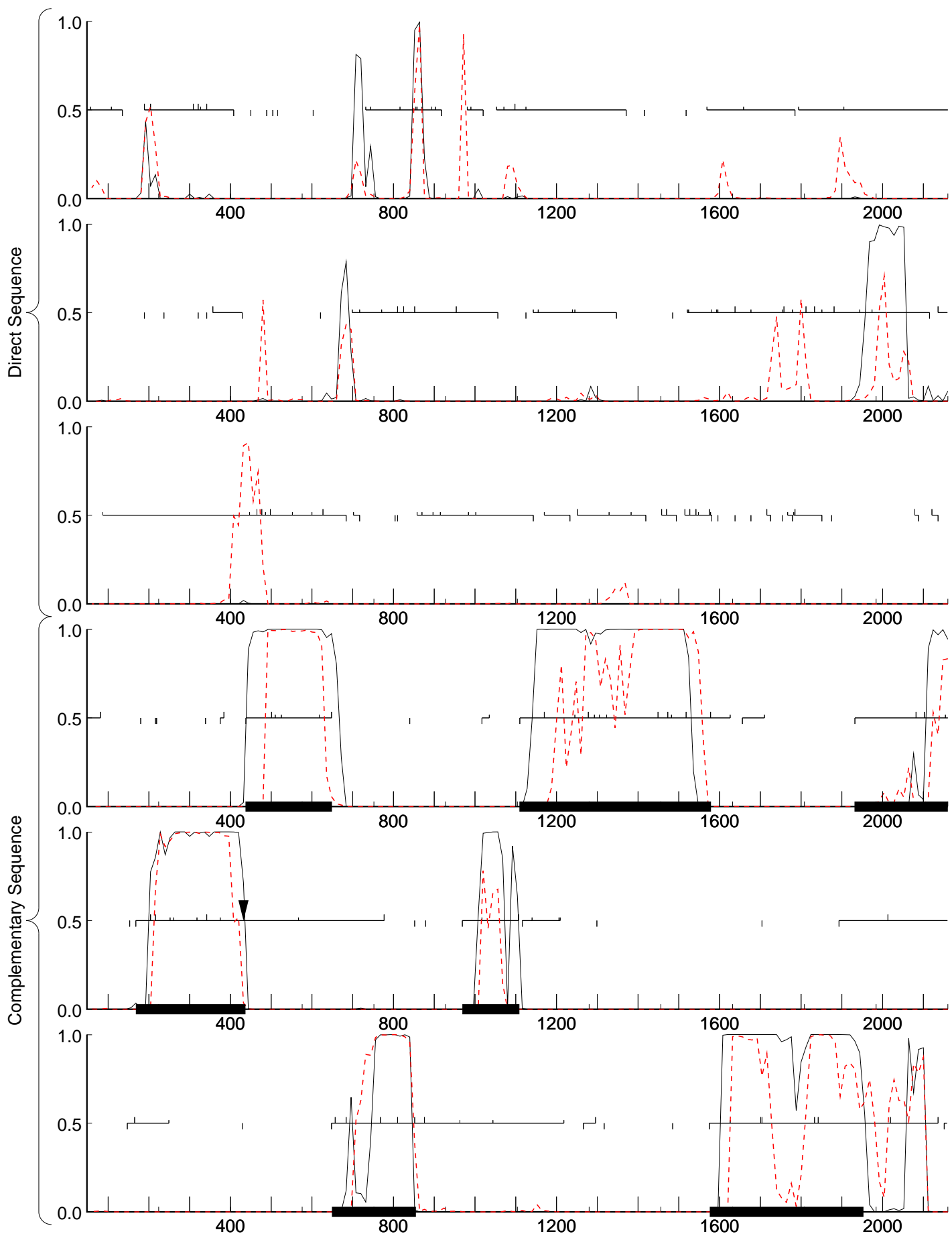
## Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014  
Sat Mar 17 16:24:50 2018

## MATRIX INFORMATION

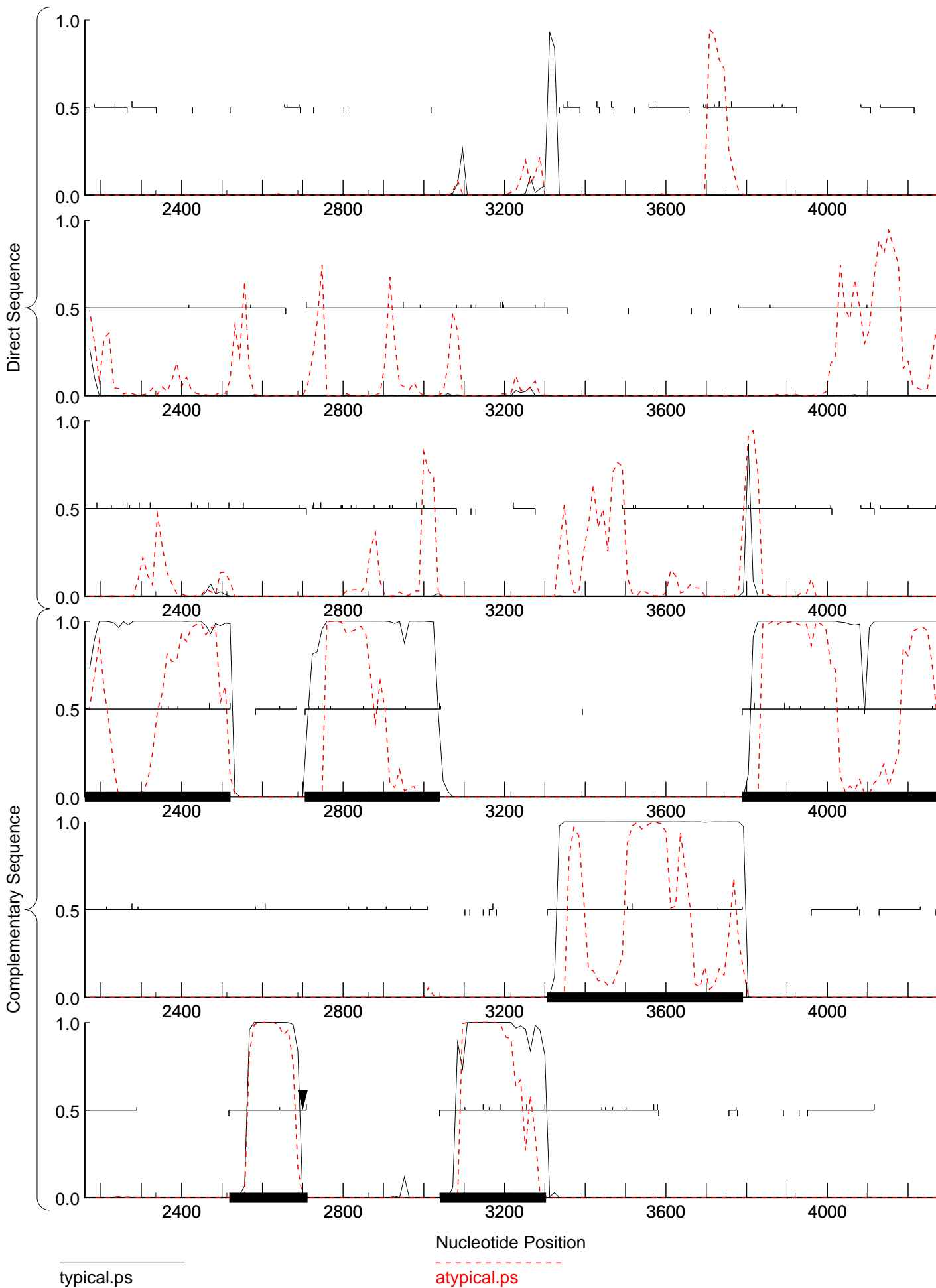
Matrix : -  
Author : -  
Order : 4

Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230



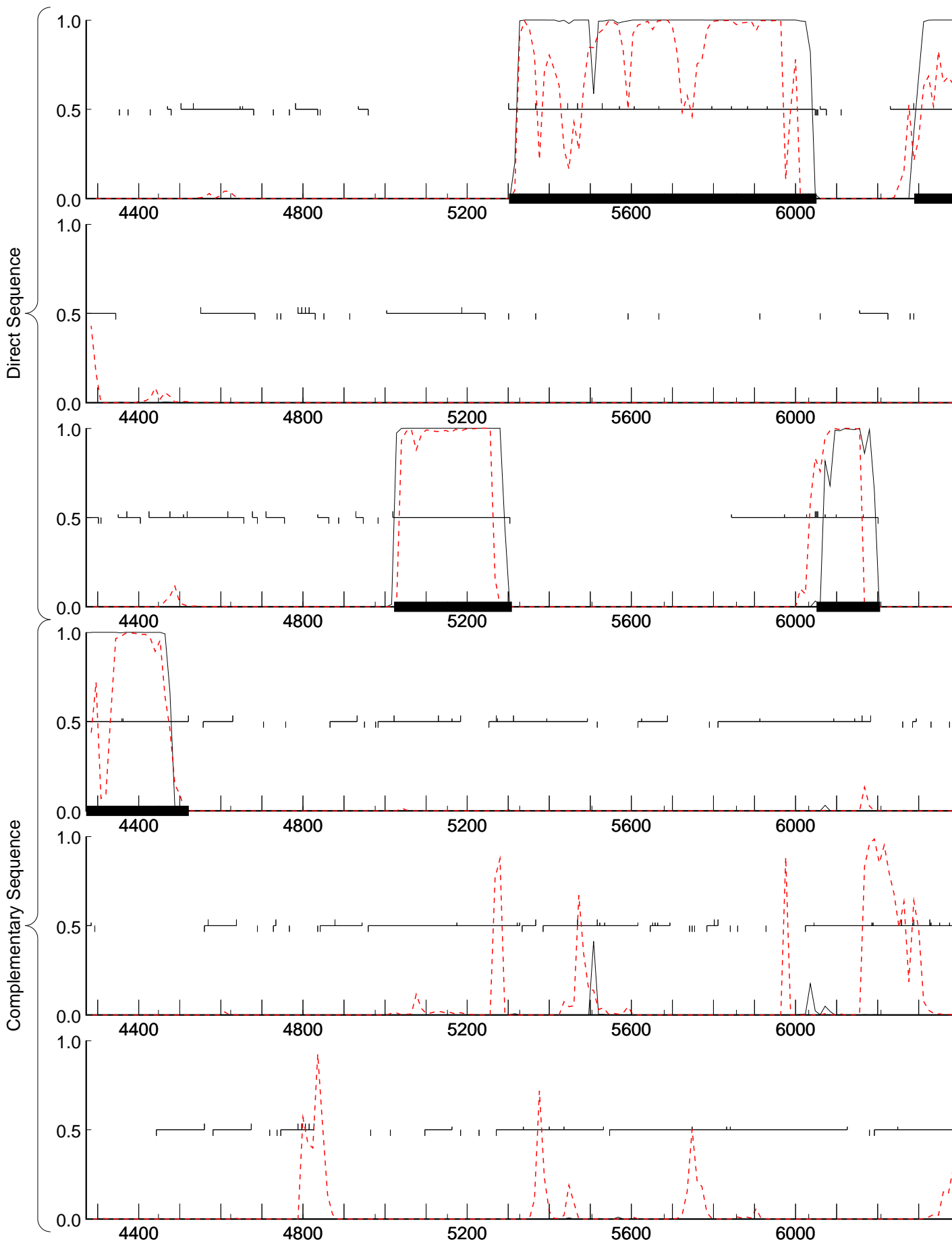
typical.ps

atypical.ps



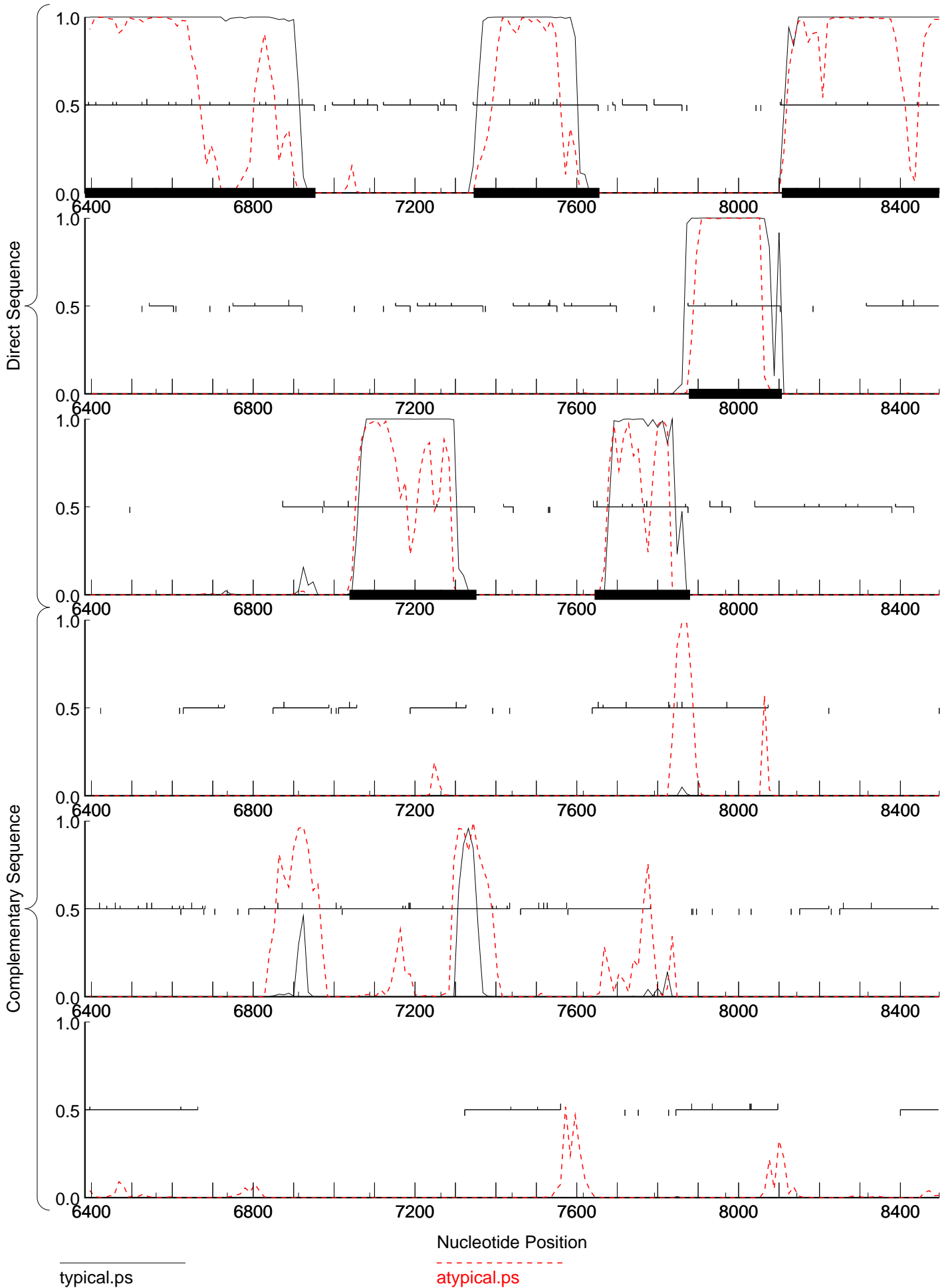
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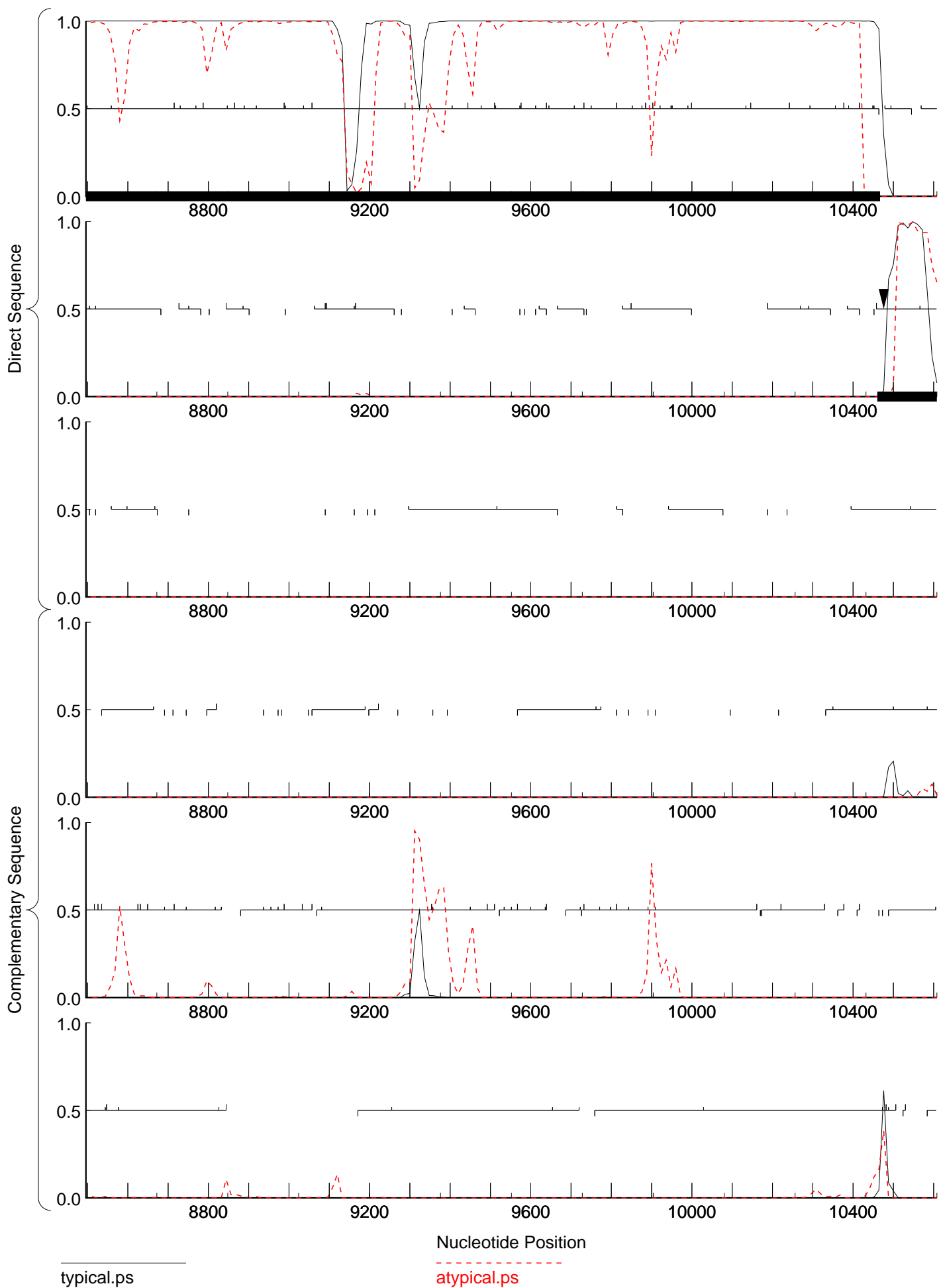
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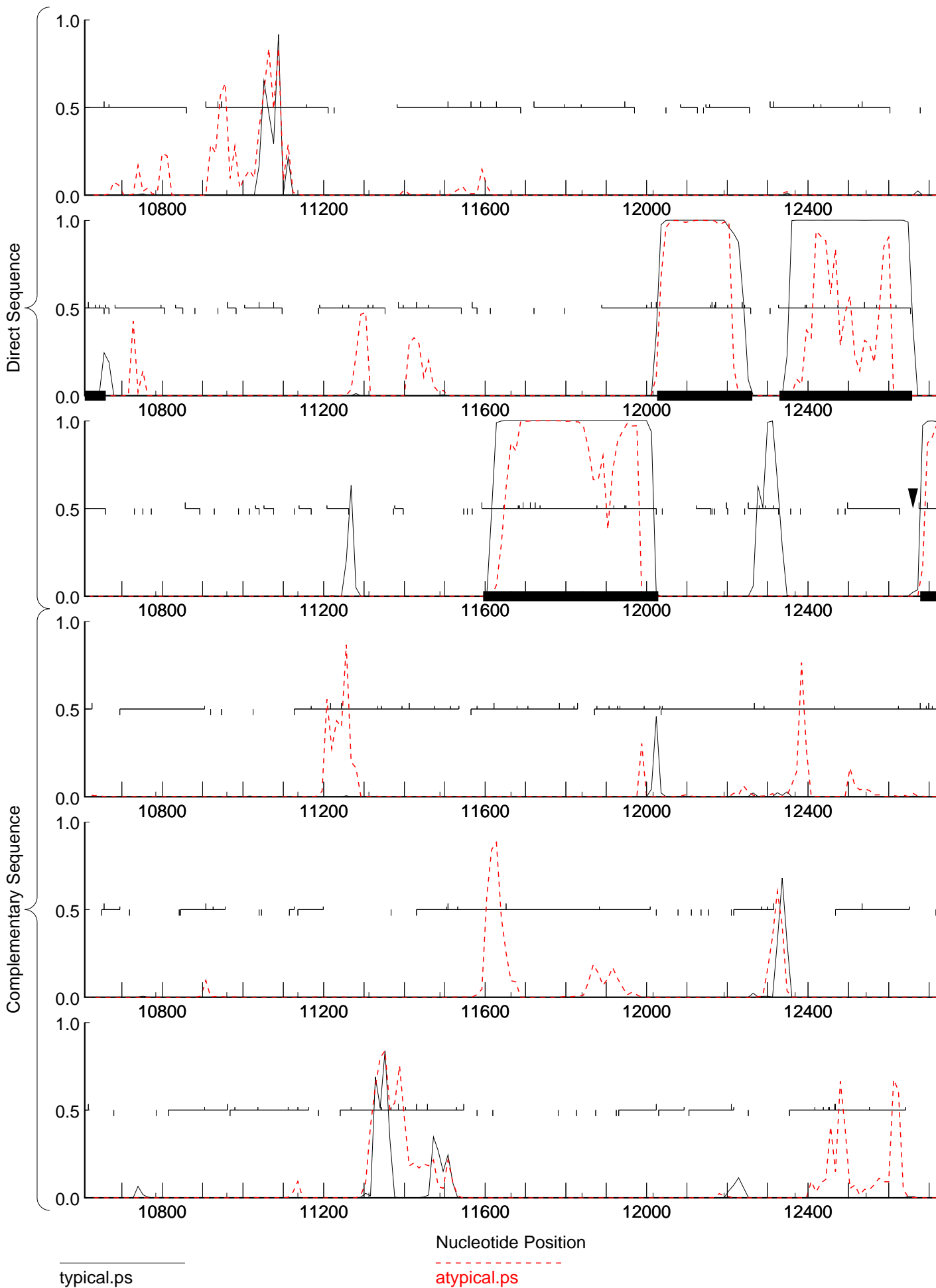


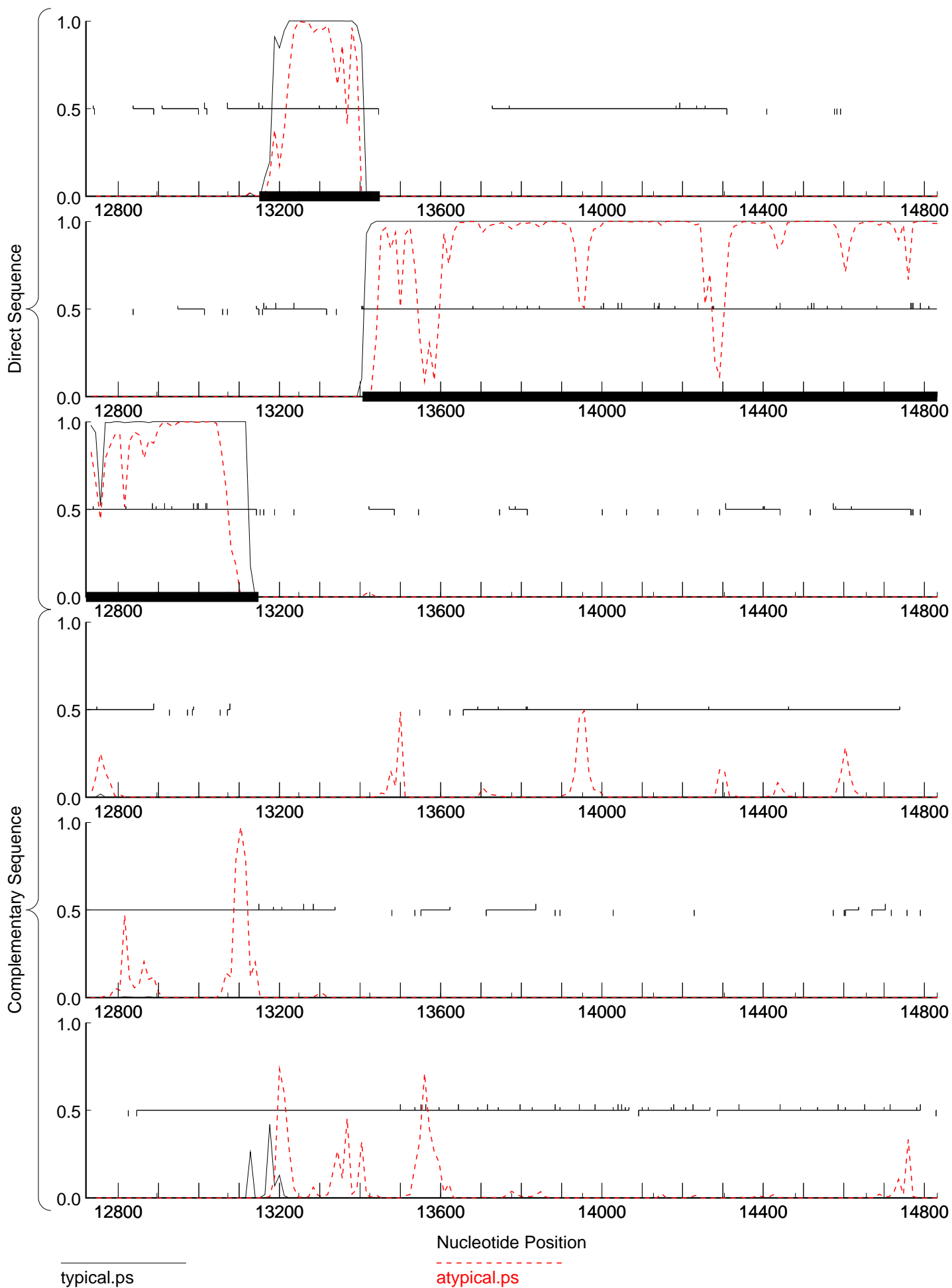
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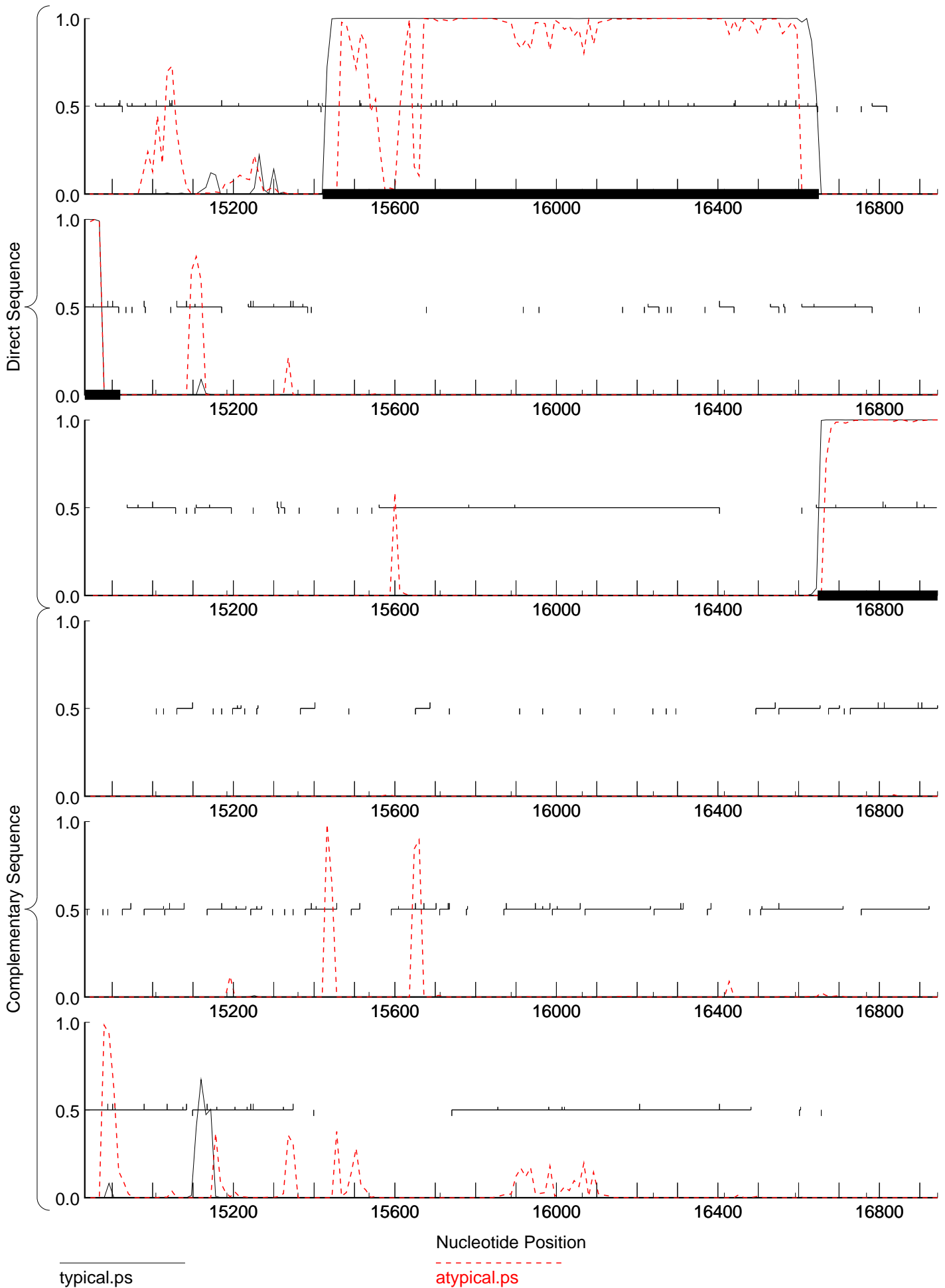


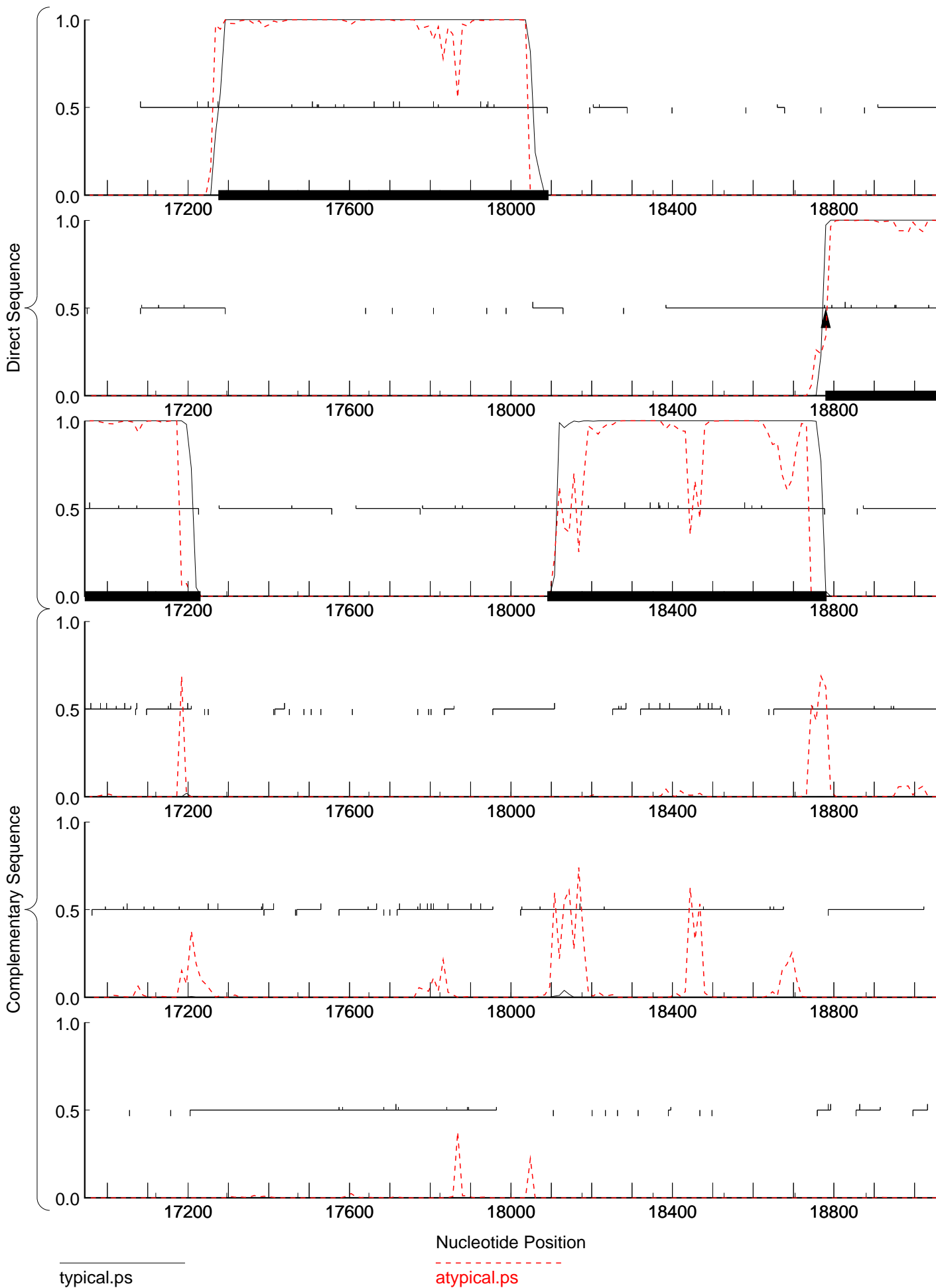






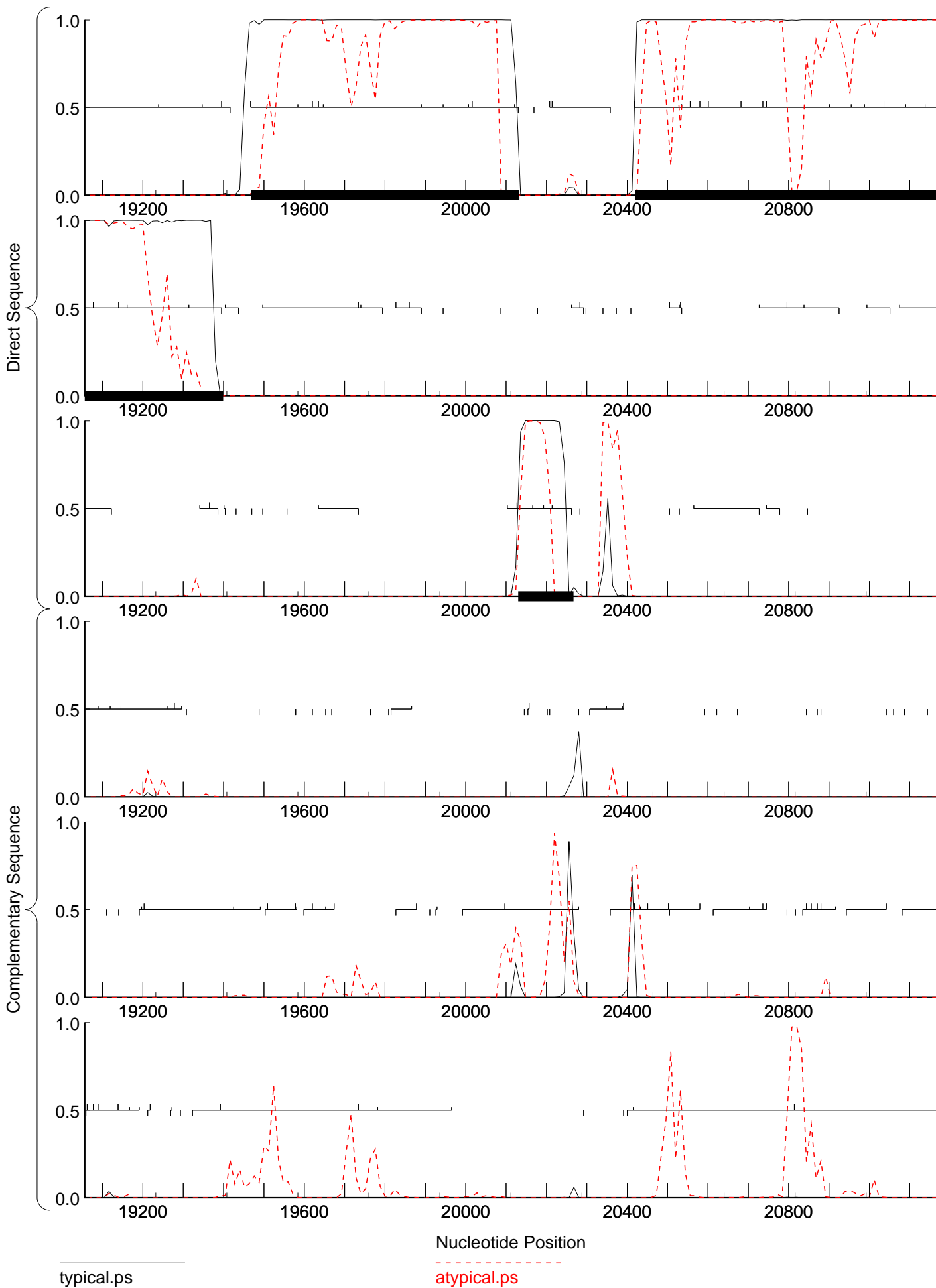


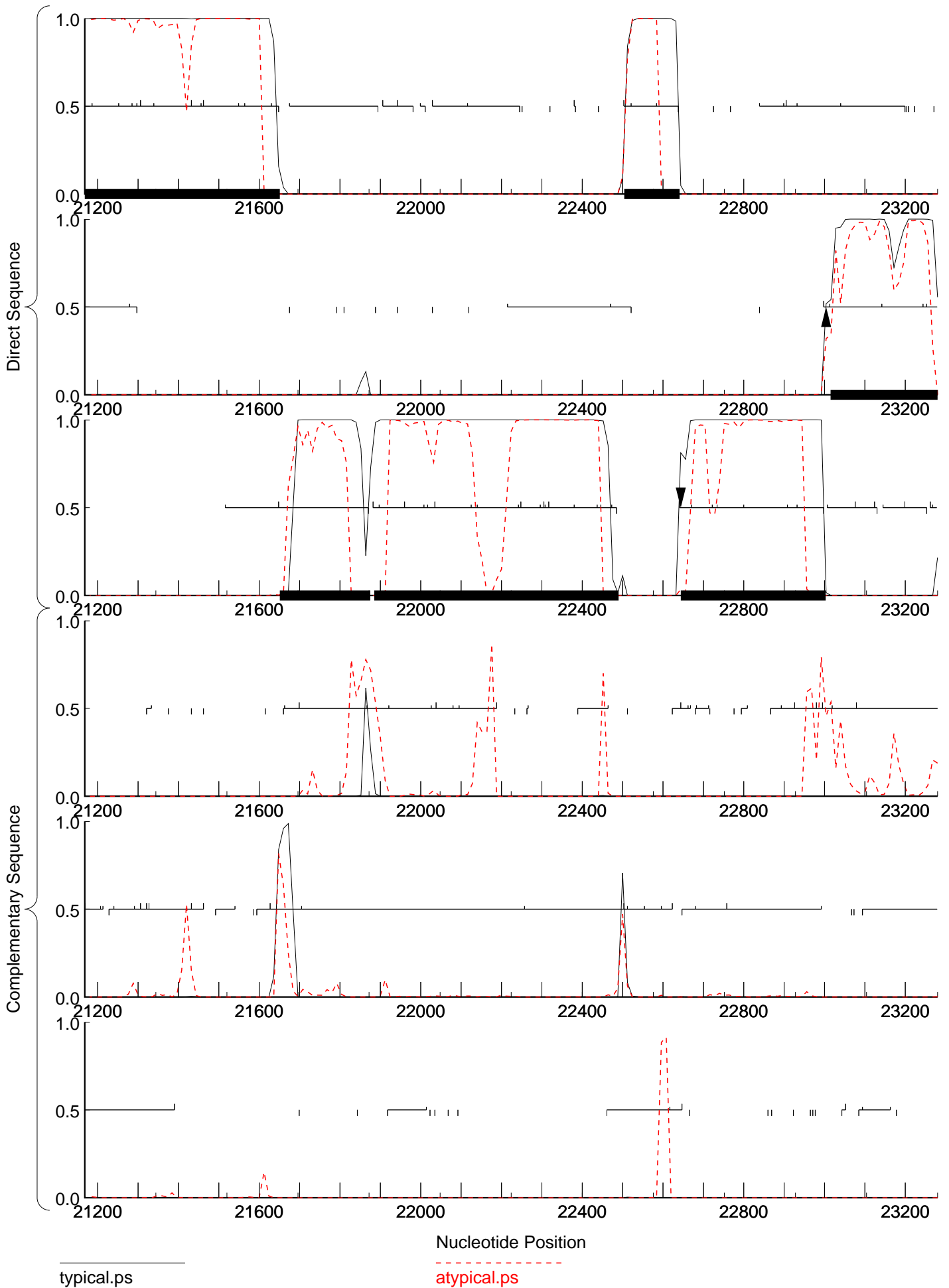


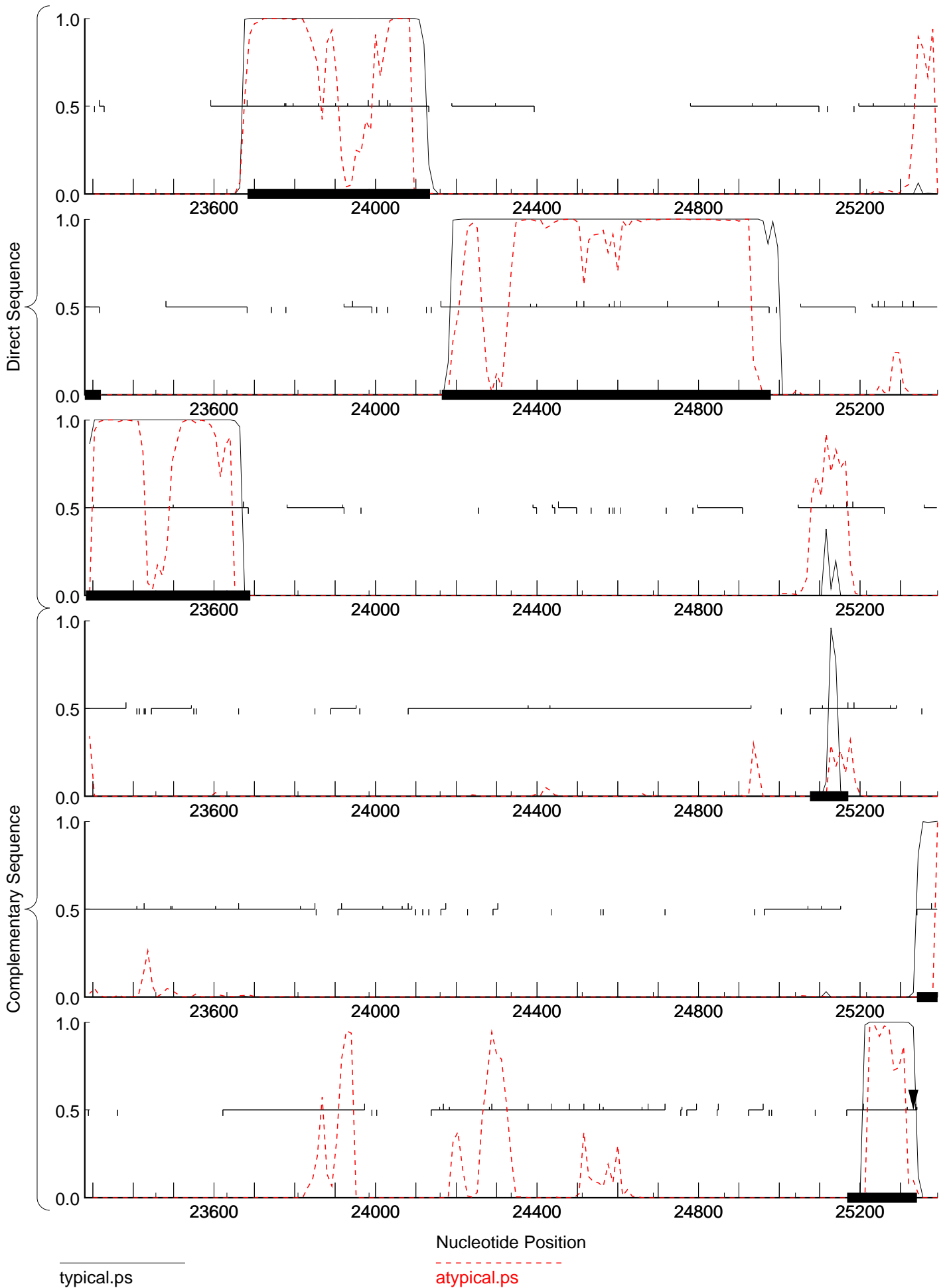


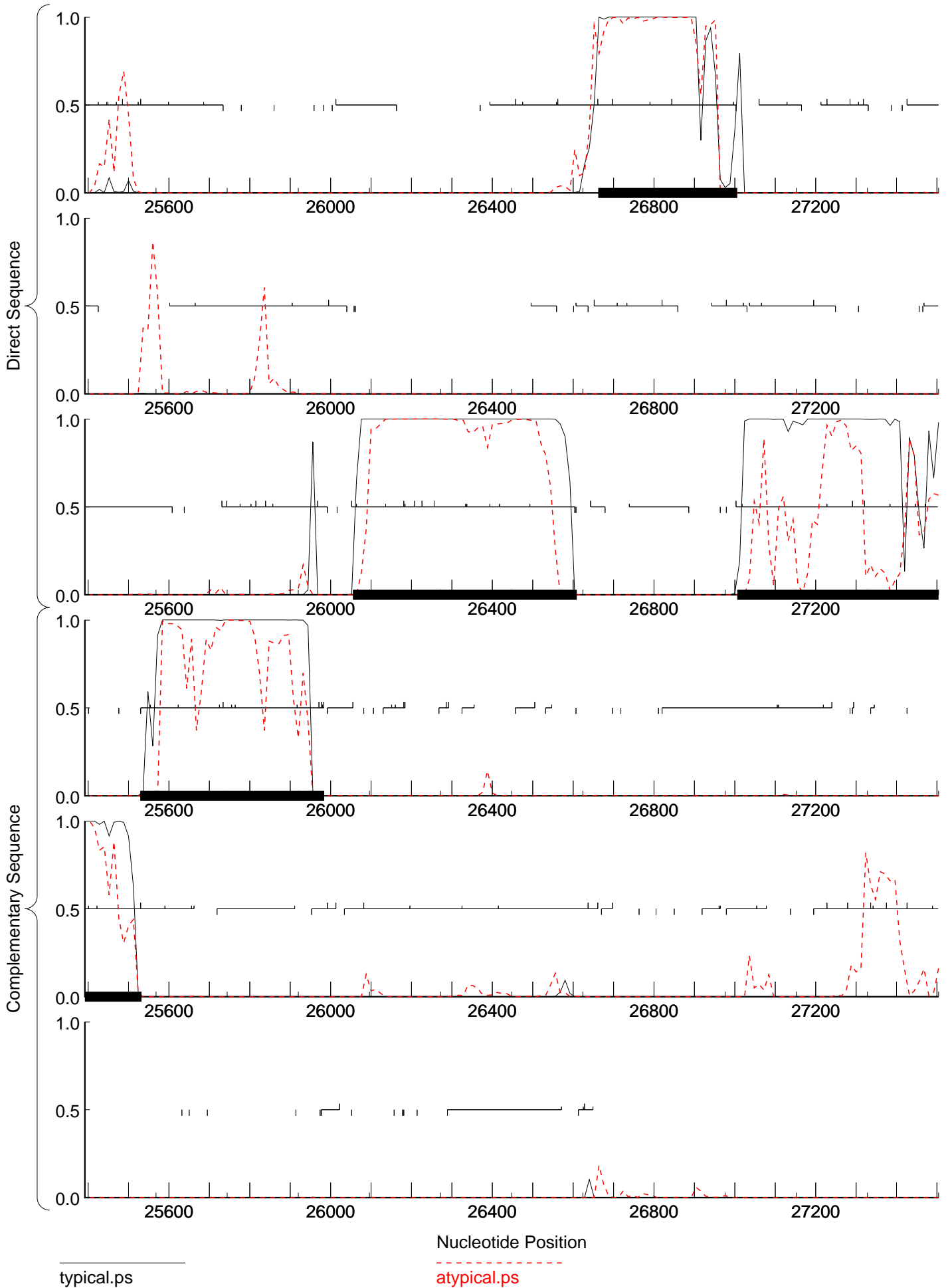
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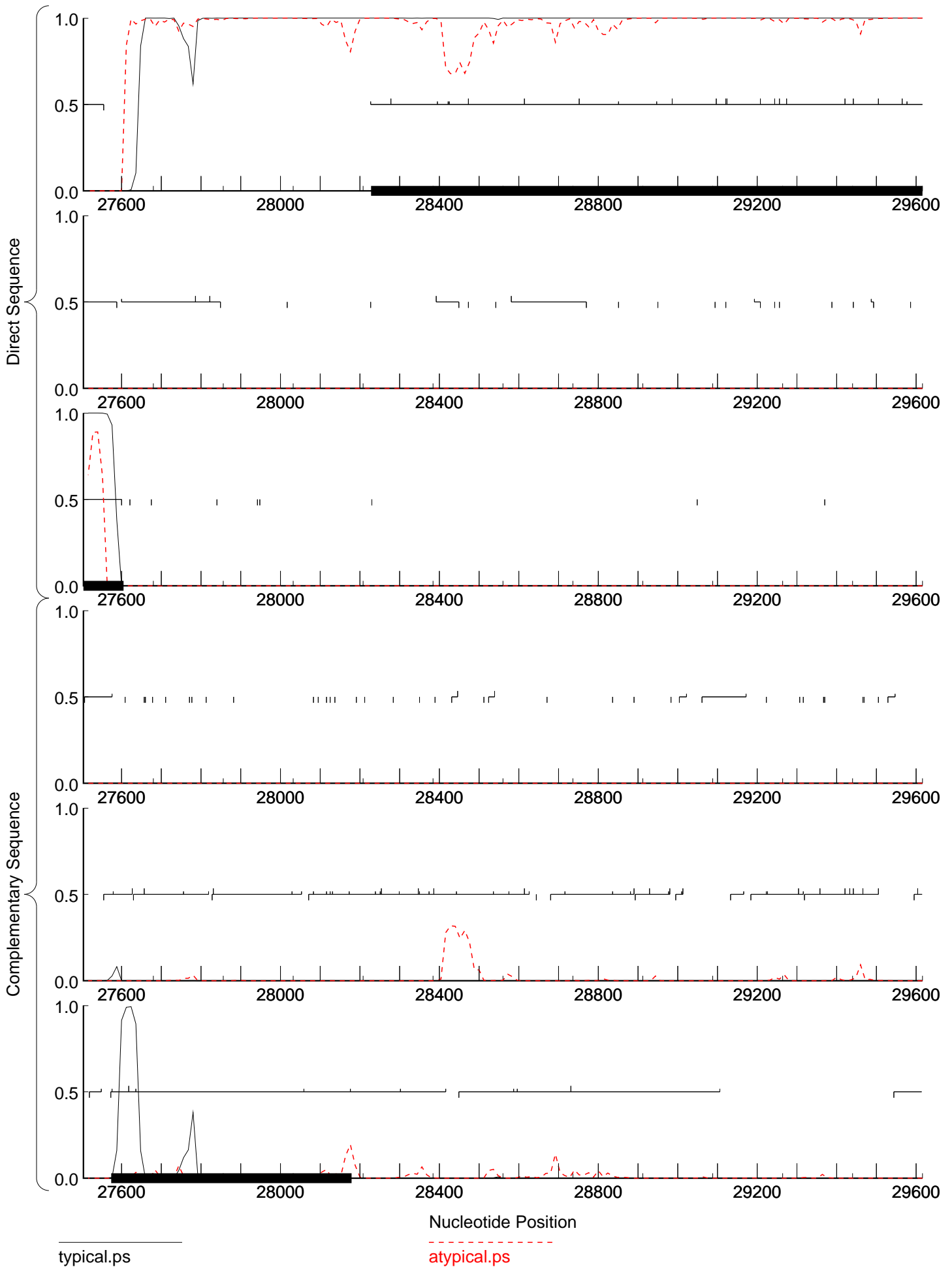
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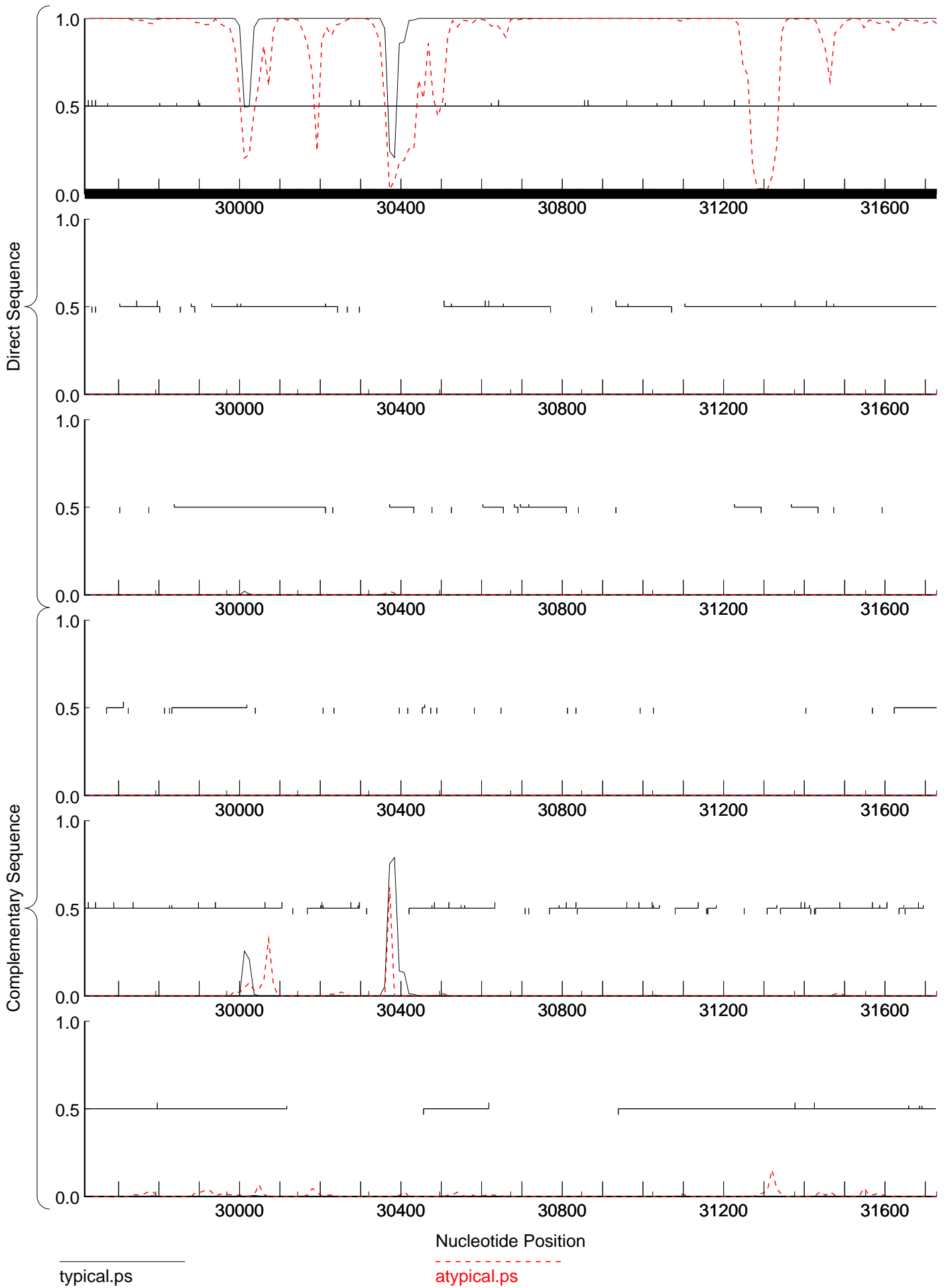




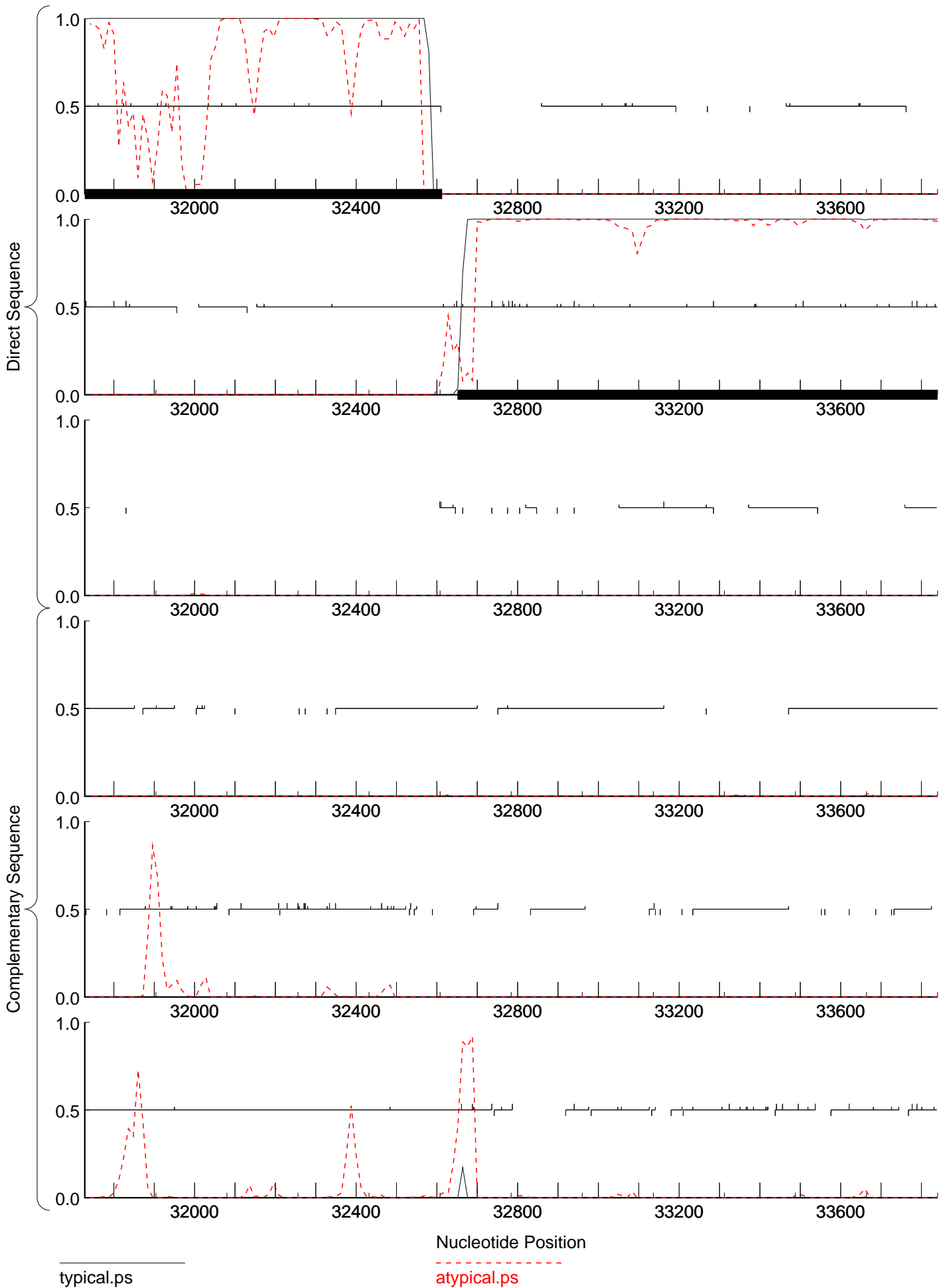


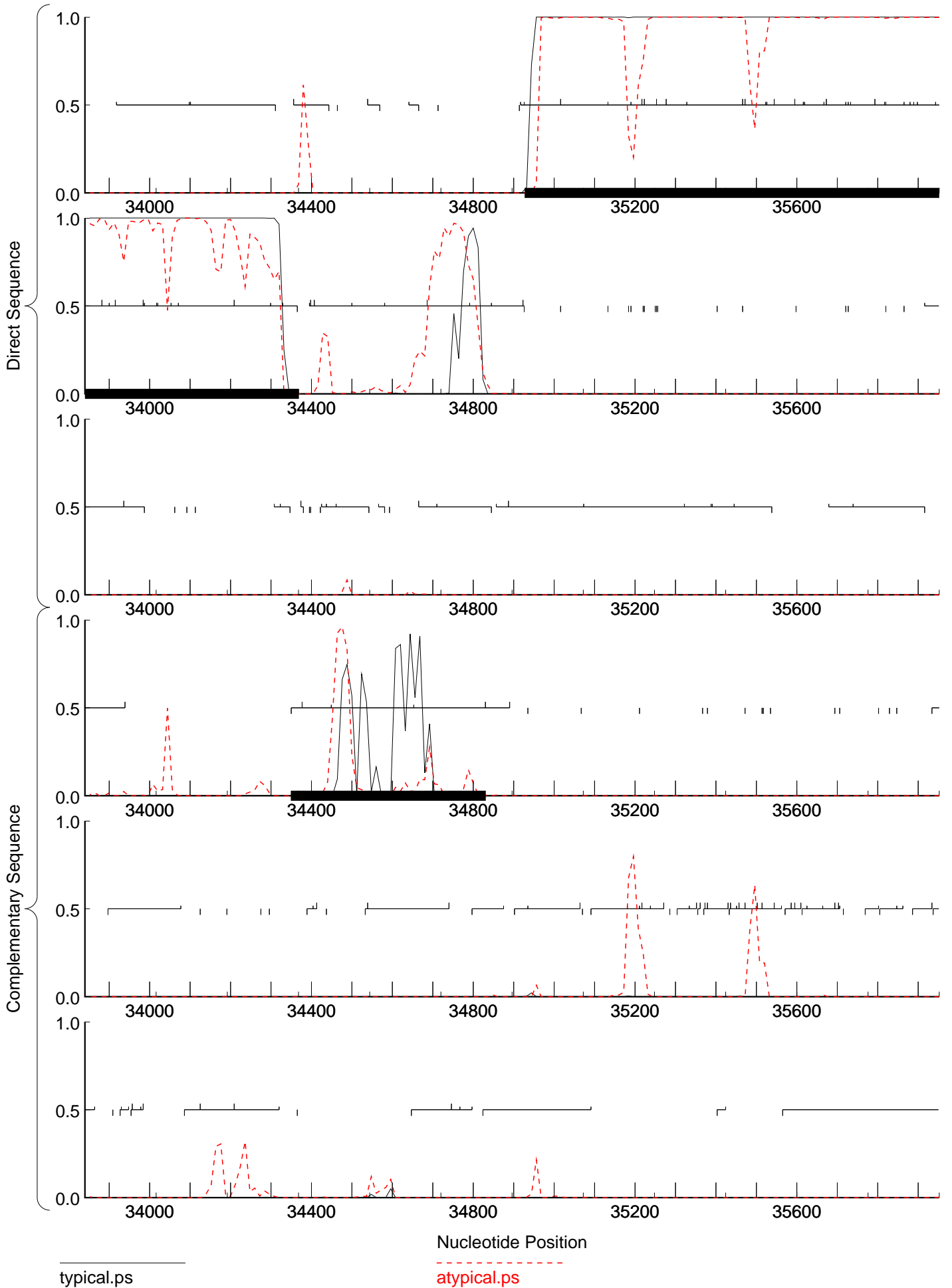
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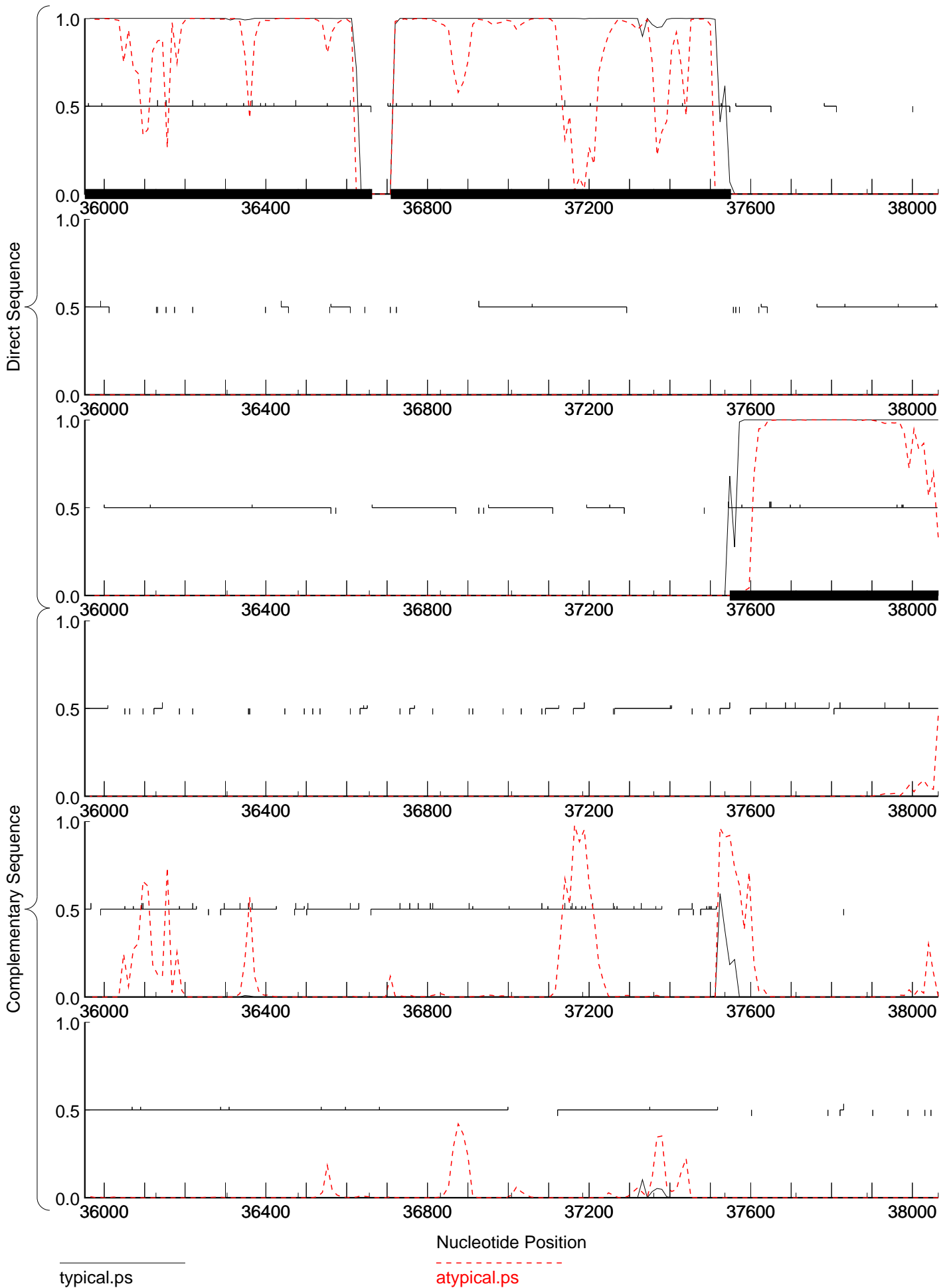
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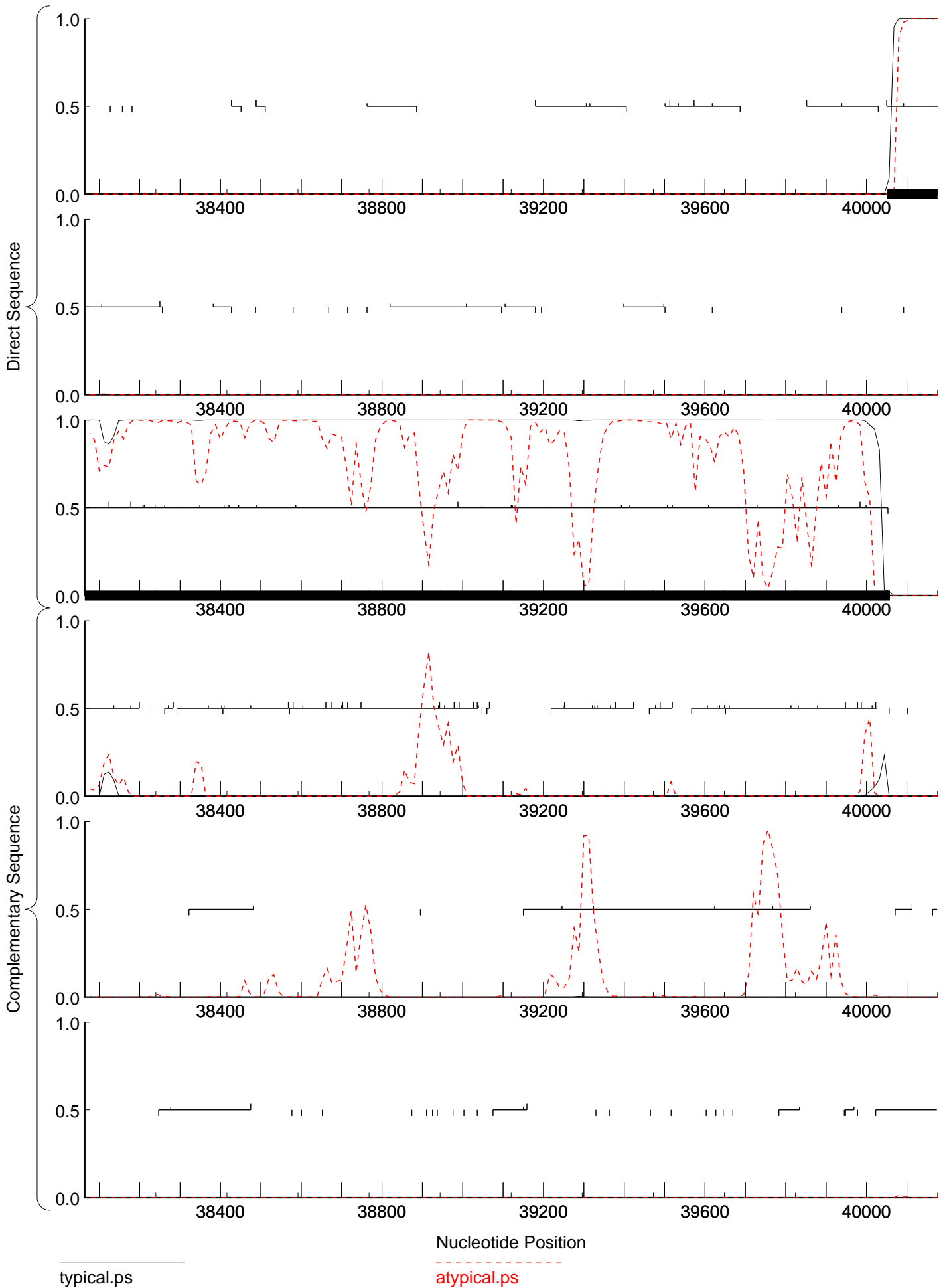


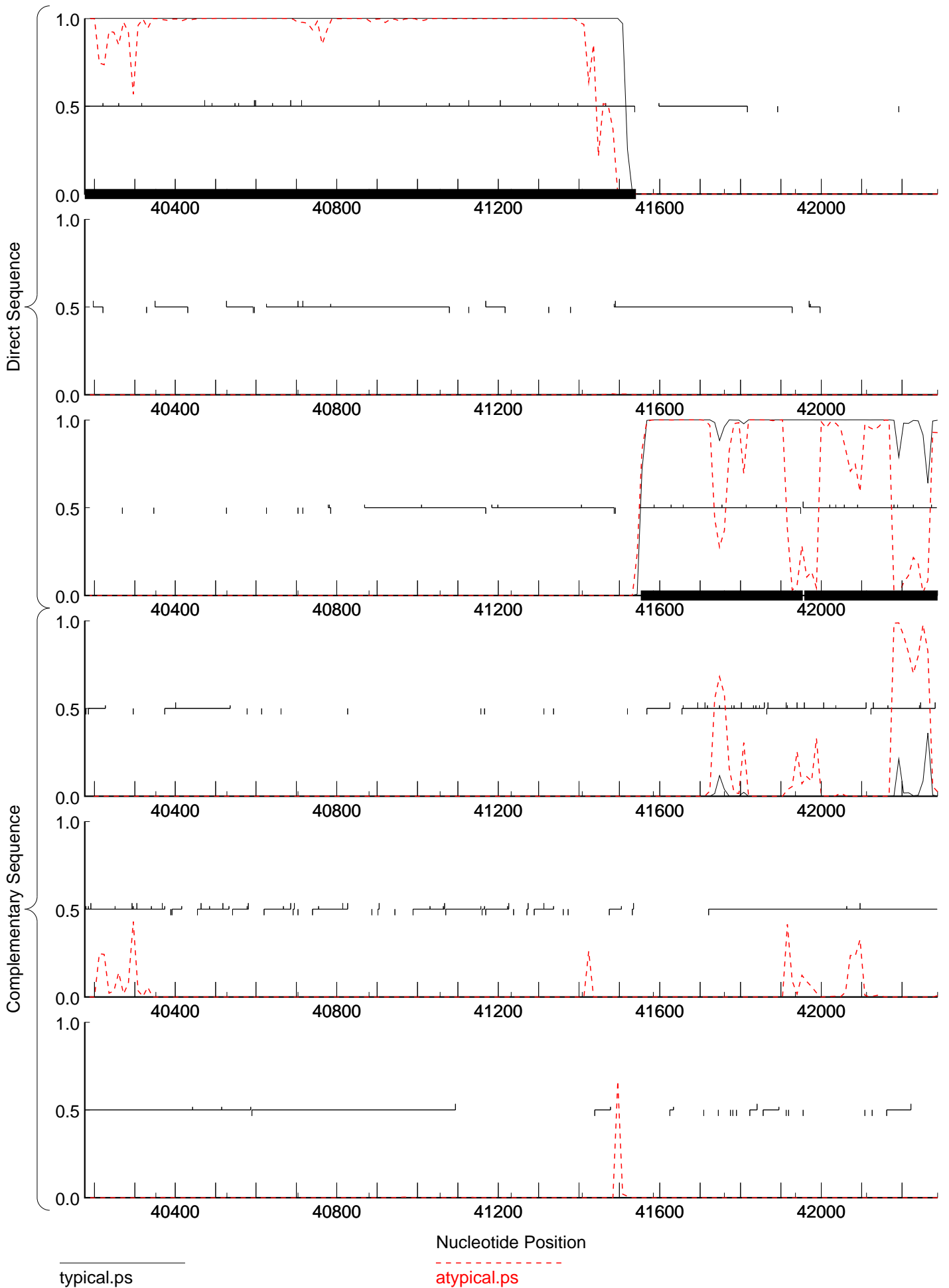


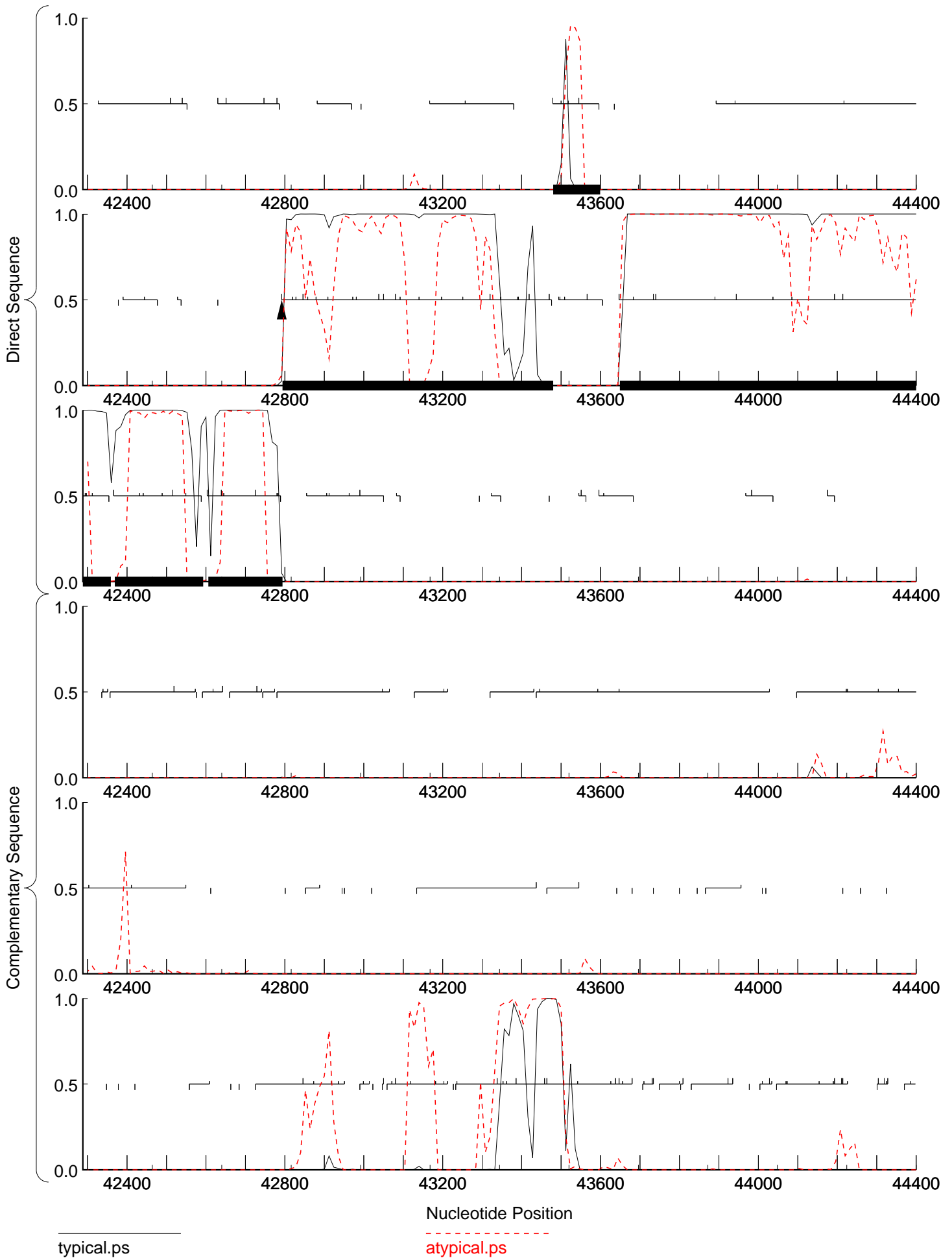


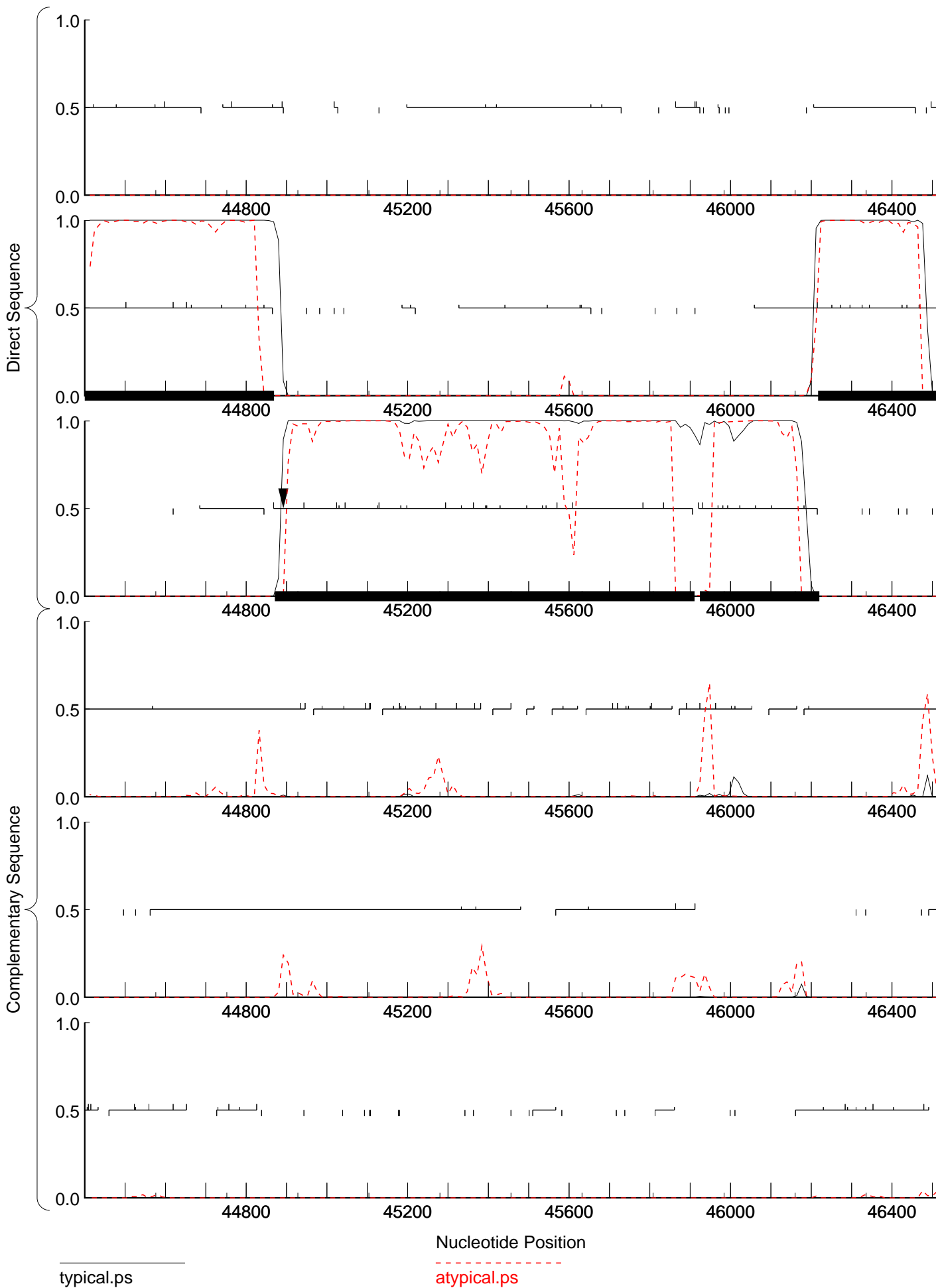


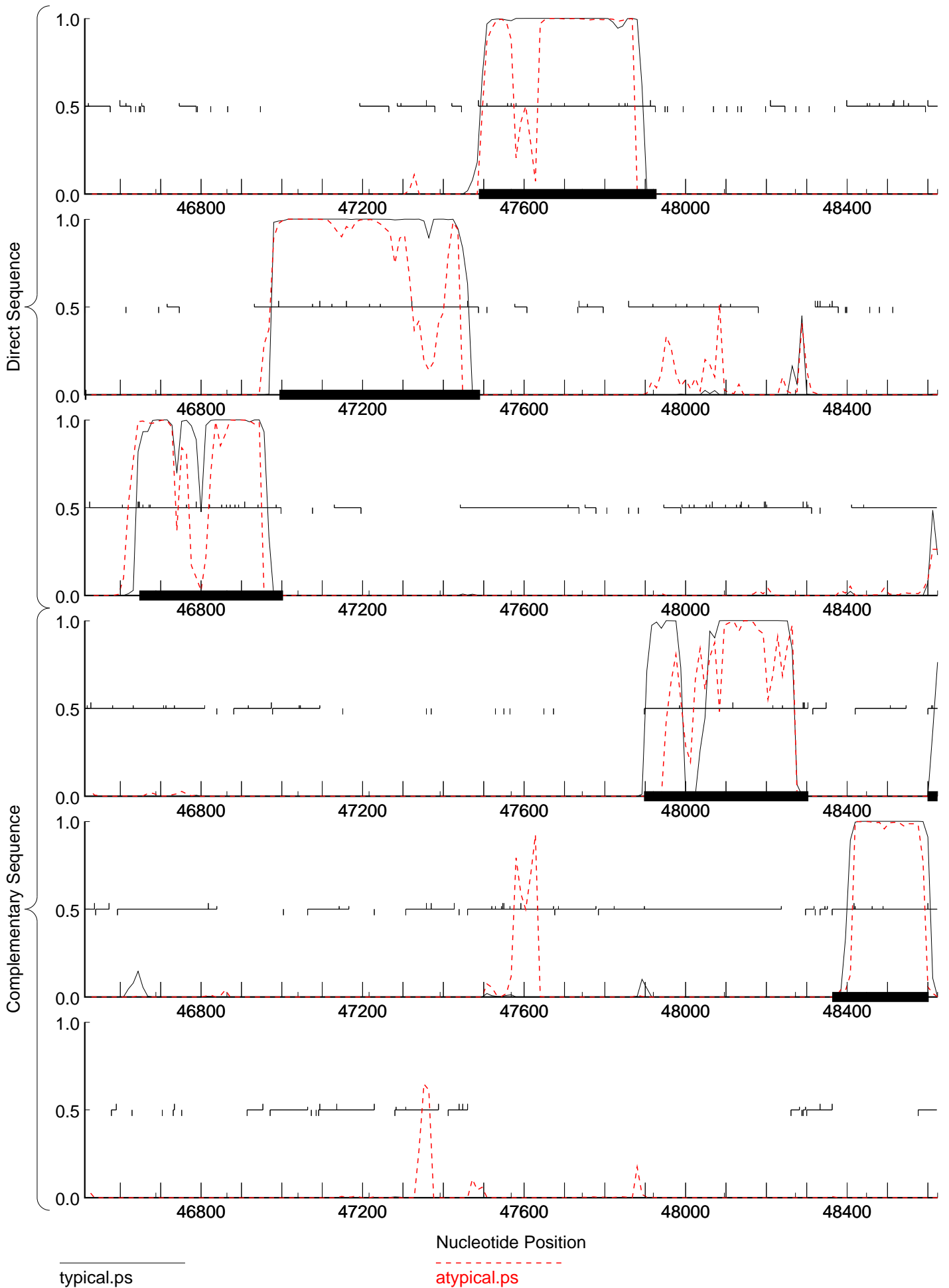








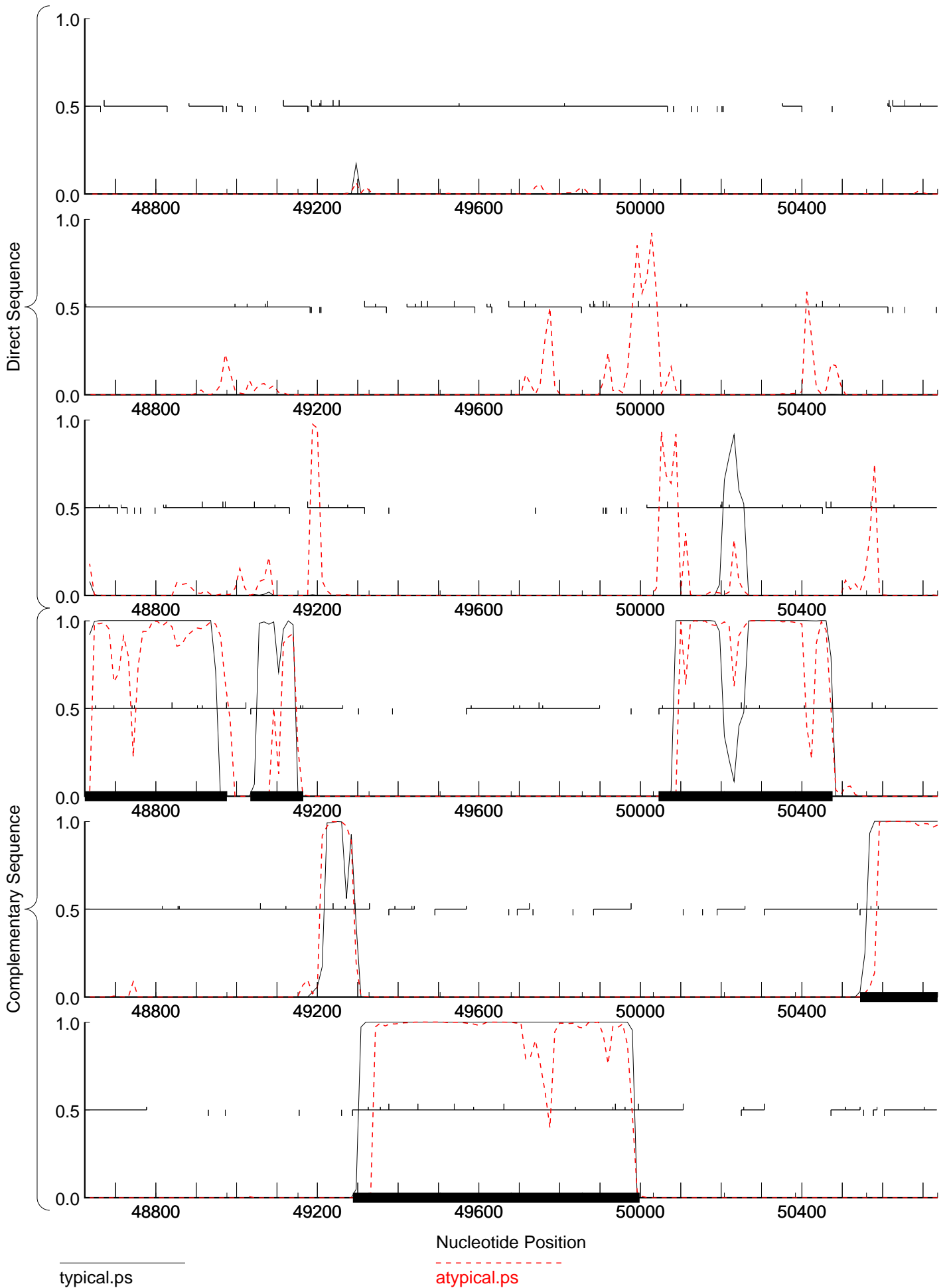


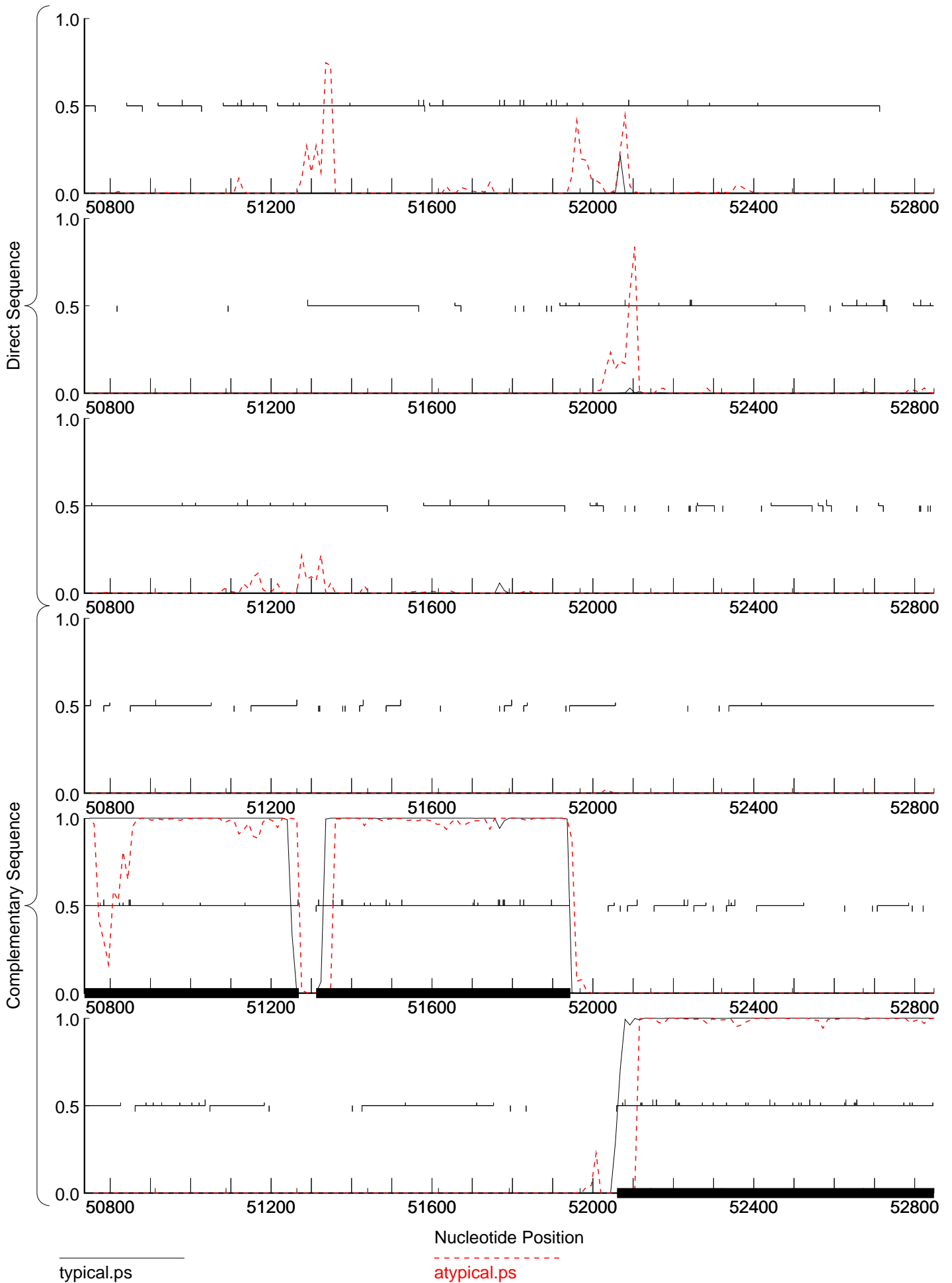


typical.ps

atypical.ps







typical.ps

atypical.ps

